

Notice of References Cited	Application/Control No. 10/591,271		Applicant(s)/Patent Under Reexamination HARBISON ET AL.	
	Examiner Ethan Whisenant		Art Unit 1634	Page 1 of 5

U.S. PATENT DOCUMENTS

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Name	Classification
*	A	US-5,686,574	11-1997	Moore et al.	530/350
*	B	US-5,989,810	11-1999	Flanagan et al.	435/6
*	C	US-6,066,452	05-2000	Weissman et al.	435/6
*	D	US-2007/0003973	01-2007	Eberwine, James H.	435/006
	E	US-			
	F	US-			
	G	US-			
	H	US-			
	I	US-			
	J	US-			
	K	US-			
	L	US-			
	M	US-			

FOREIGN PATENT DOCUMENTS

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Country	Name	Classification
	N					
	O					
	P					
	Q					
	R					
	S					
	I					

NON-PATENT DOCUMENTS

*		Include as applicable: Author, Title Date, Publisher, Edition or Volume, Pertinent Pages)
	U	Blackwell et al., Differences and similarities in DNA-binding preferences of MyoD and E2A protein complexes revealed by binding site selection. Science 250 (4984) : 1104-1110 (1990).
	V	Blackwell T.K. Selection of protein binding sites from random nucleic acid sequences. Methods in Enzymol. 254, 604-618 (1995).
	W	Bussemaker et al., Building a dictionary for genomes: identification of presumptive regulatory sites by statistical analysis. PNAS 97 (18) : 10,096-10,100 (2000).
	X	Chittenden et al.The T/E1A-Binding Domain of the Retinoblastoma Product Can Interact Selectively with a Sequence-Specific DNA-Binding Protein. Cell 65 (6) : 1073-1082 (1991).

*A copy of this reference is not being furnished with this Office action. (See MPEP § 707.05(a).)
Dates in MM-YYYY format are publication dates. Classifications may be US or foreign.

Notice of References Cited	Application/Control No. 10/591,271	Applicant(s)/Patent Under Reexamination HARBISON ET AL.	
	Examiner Ethan Whisenant	Art Unit 1634	Page 2 of 5

U.S. PATENT DOCUMENTS

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Name	Classification
	A	US-			
	B	US-			
	C	US-			
	D	US-			
	E	US-			
	F	US-			
	G	US-			
	H	US-			
	I	US-			
	J	US-			
	K	US-			
	L	US-			
	M	US-			

FOREIGN PATENT DOCUMENTS

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Country	Name	Classification
	N					
	O					
	P					
	Q					
	R					
	S					
	T					

NON-PATENT DOCUMENTS

*		Include as applicable: Author, Title Date, Publisher, Edition or Volume, Pertinent Pages)
	U	Desjarlais et al. Toward rules relating zinc finger protein sequences and DNA binding site preferences. PNAS 89(16): 7345-7349 (1992).
	V	Gelfand et al., Prediction of transcription regulatory sites in Archaea by a comparative genomic approach. Nucleic Acids Research 28(3):695-705 (2000).
	W	van Halden et al. Extracting regulatory sites from the upstream region of yeast genes by computational analysis of oligonucleotide frequencies. Journal of Molecular Biology 281 (5):. 827-842 (1998). □
	X	Kellis et al. Sequencing and comparison of yeast species to identify genes and regulatory elements. Nature 423(6937):241-54 (MAY 2003).

*A copy of this reference is not being furnished with this Office action. (See MPEP § 707.05(a).)
Dates in MM-YYYY format are publication dates. Classifications may be US or foreign.

Notice of References Cited	Application/Control No. 10/591,271		Applicant(s)/Patent Under Reexamination HARBISON ET AL.	
	Examiner Ethan Whisenant		Art Unit 1634	Page 3 of 5

U.S. PATENT DOCUMENTS

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Name	Classification
	A	US-			
	B	US-			
	C	US-			
	D	US-			
	E	US-			
	F	US-			
	G	US-			
	H	US-			
	I	US-			
	J	US-			
	K	US-			
	L	US-			
	M	US-			

FOREIGN PATENT DOCUMENTS

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Country	Name	Classification
	N					
	O					
	P					
	Q					
	R					
	S					
	T					

NON-PATENT DOCUMENTS

*		Include as applicable: Author, Title Date, Publisher, Edition or Volume, Pertinent Pages)			
	U	Kinsler et al. Whole genome PCR: application to the identification of sequences bound by gene regulatory proteins. Nucleic Acids Research 17 (10) : 3645-3653 (1989).			
	V	Liu et al., An algorithm for finding protein-DNA binding sites with applications to chromatin-immunoprecipitation microarray experiments. Nature Biotechnology 20(8):835-839 (2002).			
	W	Mavrothalassitis et al. Defining target sequences of DNA-binding proteins by random selection and PCR: determination of the GCN4 binding sequence repertoire. DNA and cell biology 9(10):783-788(1990).			
	X	McCue et al. Phylogenetic footprinting of transcription factor binding sites in proteobacterial genomes. Nucleic Acids Research 29(3) : 774-782 (2001).			

*A copy of this reference is not being furnished with this Office action. (See MPEP § 707.05(a).)
Dates in MM-YYYY format are publication dates. Classifications may be US or foreign.

Notice of References Cited	Application/Control No. 10/591,271	Applicant(s)/Patent Under Reexamination HARBISON ET AL.	
	Examiner Ethan Whisenant	Art Unit 1634	Page 4 of 5

U.S. PATENT DOCUMENTS

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Name	Classification
	A	US-			
	B	US-			
	C	US-			
	D	US-			
	E	US-			
	F	US-			
	G	US-			
	H	US-			
	I	US-			
	J	US-			
	K	US-			
	L	US-			
	M	US-			

FOREIGN PATENT DOCUMENTS

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Country	Name	Classification
	N					
	O					
	P					
	Q					
	R					
	S					
	T					

NON-PATENT DOCUMENTS

*		Include as applicable: Author, Title Date, Publisher, Edition or Volume, Pertinent Pages)
	U	McGuire et al. Conservation of DNA Regulatory Motifs and Discovery of New Motifs in Microbial Genomes. Genome Research 10 : 744-757 (2000).
	V	Nørby et al. Determination of recognition-sequences for DNA-binding proteins by a polymerase chain reaction assisted binding site selection method (BSS) using nitrocellulose immobilized DNA binding protein. Nucleic Acids Research 20(23) : 6317-6321 (1992).
	W	Pritsker et al. Whole-genome discovery of transcription factor binding sites by network-level conservation. Genome Research 14 : 99-108 (DEC 2003).
	X	Thiesen et al., Target Detection Assay (TDA): a versatile procedure to determine DNA binding sites as demonstrated on SP1 protein. Nucleic Acids Research, 18(11) : 3203-3209 (1990).

*A copy of this reference is not being furnished with this Office action. (See MPEP § 707.05(a).)
Dates in MM-YYYY format are publication dates. Classifications may be US or foreign.

Notice of References Cited	Application/Control No. 10/591,271		Applicant(s)/Patent Under Reexamination HARBISON ET AL.	
	Examiner Ethan Whisenant		Art Unit 1634	Page 5 of 5

U.S. PATENT DOCUMENTS

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Name	Classification
	A	US-			
	B	US-			
	C	US-			
	D	US-			
	E	US-			
	F	US-			
	G	US-			
	H	US-			
	I	US-			
	J	US-			
	K	US-			
	L	US-			
	M	US-			

FOREIGN PATENT DOCUMENTS

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Country	Name	Classification
	N					
	O					
	P					
	Q					
	R					
	S					
	T					

NON-PATENT DOCUMENTS

*		Include as applicable: Author, Title Date, Publisher, Edition or Volume, Pertinent Pages)
	U	Wang et al. Combining phylogenetic data with co-regulated genes to identify regulatory motifs. Bioinformatics 19(18) : 2369-2380 (DEC 2003).
	V	
	W	
	X	

*A copy of this reference is not being furnished with this Office action. (See MPEP § 707.05(a).)
Dates in MM-YYYY format are publication dates. Classifications may be US or foreign.